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0460 Page 1 of 7

0500

OIPE

RAW SEQUENCE LISTING  
PATENT APPLICATION: US/09/709,020  
DATE: 11/27/2000  
TIME: 09:23:07

Input Set : A:\ES.txt  
Output Set: N:\CRF3\11272000\I709020.raw

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3 <110> APPLICANT: Benning, Christoph  
4 Sanda, Sherrie  
5 Yu, Bin  
7 <120> TITLE OF INVENTION: Compositions and Methods for the Synthesis and Subsequent Modification of  
8 Uridine-5-Diphosphosulfoquinovose (UDP-SQ)  
10 <130> FILE REFERENCE: MSU-04769  
C--> 12 <140> CURRENT APPLICATION NUMBER: US/09/709,020  
C--> 12 <141> CURRENT FILING DATE: 2000-11-08  
12 <160> NUMBER OF SEQ ID NOS: 21  
14 <170> SOFTWARE: PatentIn version 3.0  
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17 <211> LENGTH: 1134  
18 <212> TYPE: DNA  
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28 1 5 10 15  
30 atc gtc acg cgg ctt cgg cac acg gtc gat cac ctg cag cgt ctt ggc 96  
31 Ile Val Thr Arg Leu Arg His Thr Val Asp His Leu Gln Arg Leu Gly  
32 20 25 30  
34 cac acc gtc atg gtt ttt tgc ccc gac ggc ggg ctc cgc gag cac aag 144  
35 His Thr Val Met Val Phe Cys Pro Asp Gly Gly Leu Arg Glu His Lys  
36 35 40 45  
38 ggg gct cga gtc tat ggg gtt aaa ggc ttt ccg cta ccg ctc tat ccc 192  
39 Gly Ala Arg Val Tyr Gly Val Lys Gly Phe Pro Leu Pro Leu Tyr Pro  
40 50 55 60  
42 gag ctg aag cta gct ttt ccg ttg ccg aaa gtg gga aaa gcc ttg gag 240  
43 Glu Leu Lys Leu Ala Phe Pro Leu Pro Lys Val Gly Lys Ala Leu Glu  
44 65 70 75 80  
46 cgg ttc cgg ccc gac ctg atc cac gtg gtc aat ccg gct gtg ttg ggg 288  
47 Arg Phe Arg Pro Asp Leu Ile His Val Val Asn Pro Ala Val Leu Gly  
48 85 90 95  
50 ttg ggc ggc atc tac tat gcc aag gcg cta aat gtg cca ctc gtg gcg 336  
51 Leu Gly Gly Ile Tyr Tyr Ala Lys Ala Leu Asn Val Pro Leu Val Ala  
52 100 105 110  
54 tcc tat cac acc cat ttg ccg aaa tac ctt gag cat tac ggg ctg ggg 384  
55 Ser Tyr His Thr His Leu Pro Lys Tyr Leu Glu His Tyr Gly Leu Gly  
56 115 120 125  
58 gtc ttg gag ggg gtg ctc tgg gaa ttg ctg aag ctg gcg cat aac caa 432  
59 Val Leu Glu Gly Val Leu Trp Glu Leu Leu Lys Leu Ala His Asn Gln  
60 130 135 140  
62 gca gcg atc aac ctc tgt act tca acc gcg atg gtg cag gag ctg aca 480  
63 Ala Ala Ile Asn Leu Cys Thr Ser Thr Ala Met Val Gln Glu Leu Thr

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70 gag acc ttt cgg cca gac ttg gct act gct gcg atg cgc gat cgc ctc      576
71 Glu Thr Phe Arg Pro Asp Leu Ala Thr Ala Ala Met Arg Asp Arg Leu
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74 agt ggc ggt aag ccc act gcg ccc ttg ctc tac gtc gga cgc ctc      624
75 Ser Gly Gly Lys Pro Thr Ala Pro Leu Leu Leu Tyr Val Gly Arg Leu
76          195          200          205
78 tca gcc gag aag caa atc gat cgc ctg cga ccc att ttg gat gcc aat      672
79 Ser Ala Glu Lys Gln Ile Asp Arg Leu Arg Pro Ile Leu Asp Ala Asn
80          210          215          220
82 cct gag gct tgc ttg gcc ttg gtc ggc gat gcc ccg cat cgg gcc gaa      720
83 Pro Glu Ala Cys Leu Ala Leu Val Gly Asp Gly Pro His Arg Ala Glu
84 225          230          235          240
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87 Leu Glu Gln Leu Phe Ala Gly Thr Gln Thr Gln Phe Ile Gly Tyr Leu
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94 ttt ccc tcc cgg acc gaa acc ctc ggt cta gtc ttg ctg gaa gcc atg      864
95 Phe Pro Ser Arg Thr Glu Thr Leu Gly Leu Val Leu Leu Glu Ala Met
96          275          280          285
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100          290          295          300
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106 caa ggg gcg atc gct gcg att cag cgc ttg ttg gct aac cct gca gag      1008
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108          325          330          335
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139          35          40          45
142 Gly Ala Arg Val Tyr Gly Val Lys Gly Phe Pro Leu Pro Leu Tyr Pro
143          50          55          60
146 Glu Leu Lys Leu Ala Phe Pro Leu Pro Lys Val Gly Lys Ala Leu Glu
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150 Arg Phe Arg Pro Asp Leu Ile His Val Val Asn Pro Ala Val Leu Gly
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154 Leu Gly Gly Ile Tyr Tyr Ala Lys Ala Leu Asn Val Pro Leu Val Ala
155          100          105          110
158 Ser Tyr His Thr His Leu Pro Lys Tyr Leu Glu His Tyr Gly Leu Gly
159          115          120          125
162 Val Leu Glu Gly Val Leu Trp Glu Leu Leu Lys Leu Ala His Asn Gln
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166 Ala Ala Ile Asn Leu Cys Thr Ser Thr Ala Met Val Gln Glu Leu Thr
167 145          150          155          160
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171          165          170          175
174 Glu Thr Phe Arg Pro Asp Leu Ala Thr Ala Ala Met Arg Asp Arg Leu
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178 Ser Gly Gly Lys Pro Thr Ala Pro Leu Leu Leu Tyr Val Gly Arg Leu
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182 Ser Ala Glu Lys Gln Ile Asp Arg Leu Arg Pro Ile Leu Asp Ala Asn
183          210          215          220
186 Pro Glu Ala Cys Leu Ala Leu Val Gly Asp Gly Pro His Arg Ala Glu
187 225          230          235          240
190 Leu Glu Gln Leu Phe Ala Gly Thr Gln Thr Gln Phe Ile Gly Tyr Leu
191          245          250          255
194 His Gly Glu Gln Leu Gly Ala Ala Tyr Ala Ser Ala Asp Ala Phe Val
195          260          265          270
198 Phe Pro Ser Arg Thr Glu Thr Leu Gly Leu Val Leu Leu Glu Ala Met
199          275          280          285
202 Ala Ala Gly Cys Pro Val Val Ala Ala Asn Ser Gly Gly Ile Pro Asp
203          290          295          300
206 Ile Val Ser Asp Gly Ile Asn Gly Phe Leu Phe Asp Pro Glu Asp Glu
207 305          310          315          320
210 Gln Gly Ala Ile Ala Ala Ile Gln Arg Leu Leu Ala Asn Pro Ala Glu
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214 Arg Glu Ile Leu Arg Gln Ala Ala Arg Gln Glu Ala Glu Arg Trp Ser
215          340          345          350
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DATE: 11/27/2000

PATENT APPLICATION: US/09/709,020

TIME: 09:23:07

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Output Set: N:\CRF3\11272000\I709020.raw

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242 aaccacaagc tcaatatagg ttgaccata agaacaatca atgcaagatc attttgtgta 360
244 ccagtcctatg attgaataaa gtttcagttc ggttacagct cgtttataag aaaattggca 420
246 gaaattgttt ttcaaccat ttccgttcgg ttgatagct catcaatatg gtttggcagt 480
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250 ggcacgggcc gcgacgacag gaagtcctga agacattgct tcaagaacca caaagccaa 600
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RAW SEQUENCE LISTING  
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